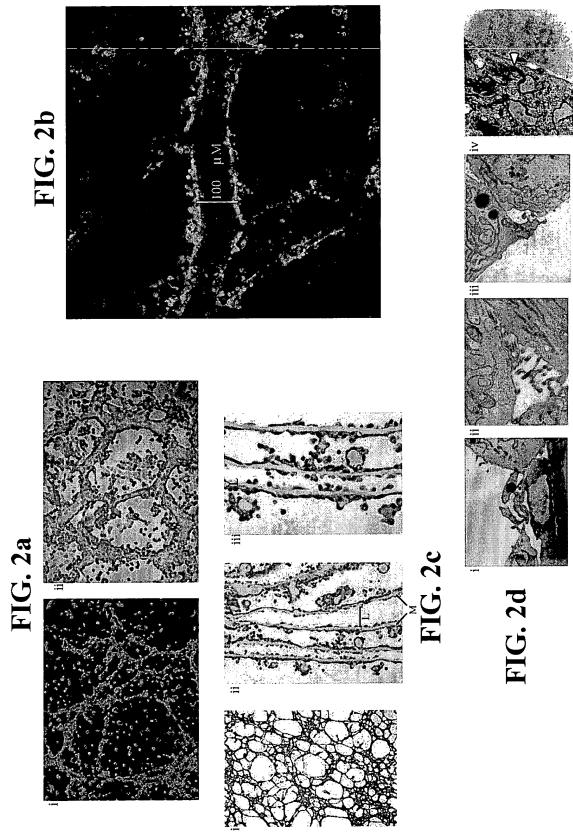


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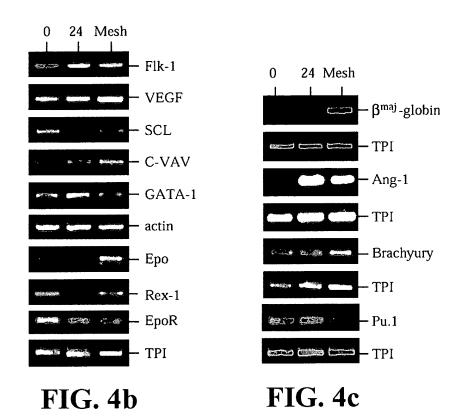


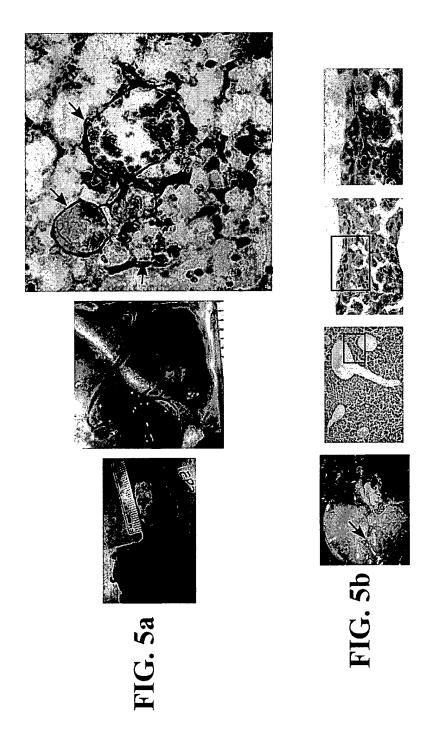


3/10 Tien Thy. - scal Predectin CD31 CDA5 CD3A FIG. 3a 🕨

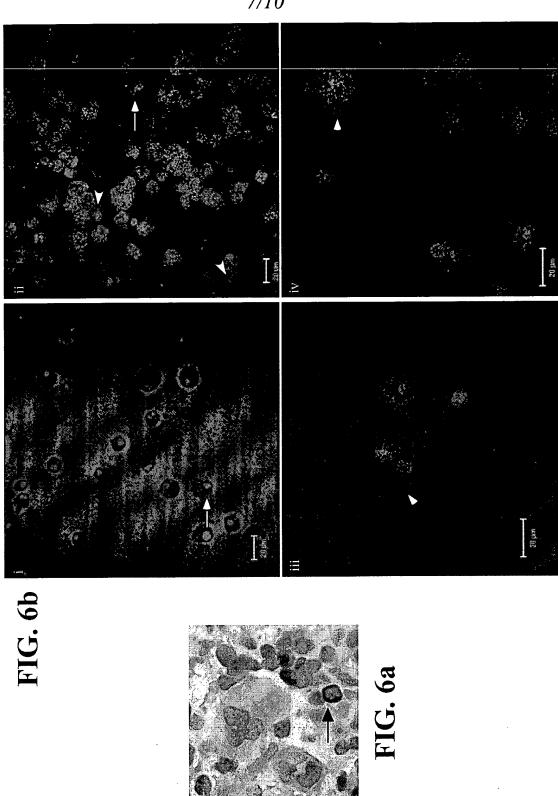
4/10																							
Lower Primer	CGTCGTATTCCTGTTTGCTG	GCAACAACGGAGGACATTAG	GGAGGACTGGAGAAATCAG	TGCTAGCCAATTCCTCCCAG	GATGAGGACCAGAATGAGAGAC	CATCCGGAACAATCTCTTTTC	AGTGTCTGTAGGCCTCAGCT	CAGCCATCAAAAGGACACAC	CATACAGTACGACACTGACG	GATGGCCAGTGCTCATTGTTT	GTTCTGTCTTTCTTTGGTCTG		Lower Primer	ACTTGCTGCTGCAACGGAGAC	TGAAATCAGCACCGTGTAAG	TGCTGCCTGTGAGTCATAAC	GCAACAACGGAGGACATTAG	AAT GAG GGG ATG GGG AGG AAG	CCT CGC CTG TCT TGC CGT AGT T	GTCCCCATGGAGTCAAAGAG	CTCAAGGAGCTTTGCTCA	CACAACCCCAGAAACAGACA	CAGGCAAGCAGGAAAGGGG
	SEQ ID NO: 4	SEQ ID NO: 6	SEQ ID NO: 8	SEQ ID NO: 10	SEQ ID NO: 12	SEQ ID NO: 14	SEQ ID NO: 16	SEQ ID NO: 18	SEQ ID NO: 20	SEQ ID NO: 22	SEQ ID NO: 24			SEQ ID NO: 26	SEQ ID NO: 28	SEQ ID NO: 30	SEQ ID NO: 32	SEQ ID NO: 34	SEQ ID NO: 36	SEQ ID NO: 38	SEQ ID NO: 40	SEQ ID NO: 42	SEQ ID NO: 44
Upper Primer	CCGCCTCCTCCTCCTG	GAAGCCAAGGACAGACAC	GTATAACGTGGAGGTCAAGC	ATGTCACTGCTGGTGCTGGA	CCTATGACCACCACATCCG	CGCTCTGTGGTTCTGCGTG	ACT CGT CAT ACC ACT AAG GT	ACTGACCCTAAAGCAAGACG	ATTGCACACACGGGATTCTG	CCCTGGCATGATCAAAGACTT	TGAACTTTCTGCTCTTTGGG		Upper Primer	GGGAGGAAAAAGAGAG	GGGAGGAAAAGAGAG	TCCAGGTGCTATATTGCC	GAAGCCAAGGACAGACAC	GAG ACA GGC AGC AAG AAA AAG	GGA CAG CAT CTG GTG GGT GGA C	AAGGAAAGAAGGAAAG	GTCCCCATGGAGTCAAAGA	CTGACAGATGCTCTTTGGG	TGACTGATGCTGAGAAGGCTG
	SEQ ID NO: 3	SEQ ID NO: 5	SEQ ID NO: 7	SEQ ID NO: 9	SEQ ID NO: 11	SEQ ID NO: 13	SEQ ID NO: 15	SEQ ID NO: 17	SEQ ID NO: 19	SEQ ID NO: 21	SEQ ID NO: 23			SEQ ID	SEQ ID NO: 27	SEQ ID NO: 29	SEQ ID NO: 31	SEQ ID NO: 33	SEQ ID	SEQ ID NO: 37	SEQ ID	SEQ ID	SEQ ID
<u>RT-PCR</u> <u>Gene</u> <u>Siz</u> e	Actin 402	Brachyury 194	C-VAV 604	Epo 409	EpoR 702	FIK-1 500	GATA-1 263	Rex-1 270	SCL 744	TPI 523	VEGF 406	Nesteled PCR	Gene Size	Ang-1 770	458	Brachyury 947	194	PU.1 648	164	βH1 globin 792	265	β ^{ma} globin 578	457

FIG. 4a





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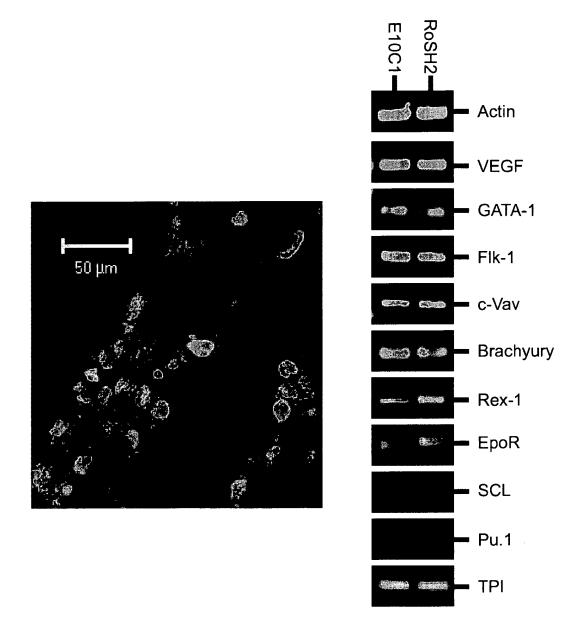


FIG. 7

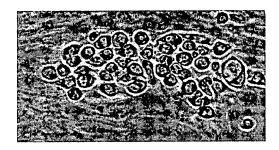
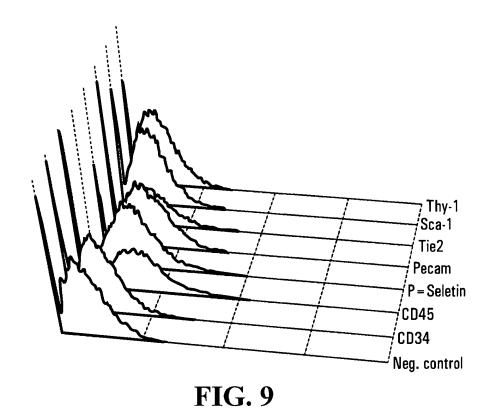


FIG. 8



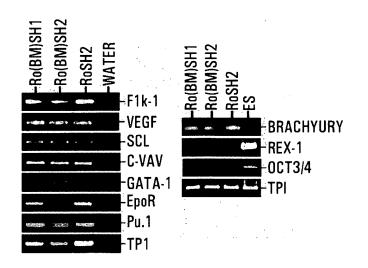


FIG. 10